

Applications using

Analysis of categorical variables

Define the categorical variables *cut* and *color*

```
cut=diamonds$cut  
color=diamonds$color
```

Display the categories of the categorical variable *cut*

```
unique(cut)
```

Display the absolute frequencies of the categorical variable *cut* and identify the mode

```
table(cut)
```

Draw a bar plot of the categorical variable *cut*

```
plot(cut)
```

or

```
a=ggplot(diamonds, aes(cut))  
b=a+geom_bar()  
b
```

Display the percentages of the categorical variable *cut*

```
100*table(cut)/length(cut)
```

or

```
100*prop.table(table(cut))
```

Draw the pie chart of the variable *cut*

```
pie(table(cut))
```

Draw a bar plot of the categorical variables *cut* and *color*

```
plot(cut,color)
```

or

```
a=ggplot(diamonds, aes(cut,color))  
b=a+geom_count()
```

Run the chi-square test to study the association between two categorical variables, *cut* and *color*

Null hypothesis $H_0: \chi^2 = 0$ (independence, no association)

Alternative hypothesis $H_1: \chi^2 > 0$ (dependence, association)

The decision of the test depends on the *p*-value.

If *p*-value < 0.05, there is an evidence against H_0 (we reject H_0).

If *p*-value ≥ 0.05 , there is an evidence in favor of H_0 (we do not reject H_0)

```
chisq.test(cut,color)
```

Analysis of numerical variables

Install and load the package *ggplot2*

Consider the dataset *diamonds*

View the dataset *diamonds*

`View(diamonds)`

Define the variable *carat* from the dataset *diamonds*

`carat=diamonds$carat`

Compute the mean, the median, the minimum, and the maximum of the variable *carat*

`m=mean(carat)`

`Me=median(carat)`

`min(carat)`

`max(carat)`

Compute the first and third quartile of the variable *carat*

`Q1=quantile(carat,0.25)`

`Q3=quantile(carat,0.75)`

Compute the p-quantile of the variable *carat*

`Qp=quantile(carat,p)`

Compute the frequency of the value 0.21 for the variable *carat*

`length(carat[carat==0.21])`

Compute minimum value, maximum value, mean, median, first and third quartile of the variable *carat*

`summary(carat)`

Draw the histogram of the variable *carat*

`hist(carat)`

or

`a=ggplot(diamonds, aes(x=carat))`

`b=a+geom_histogram()`

`b`

Draw the histogram of the variable *carat* with 50 bins

`hist(carat,50)`

or

`a=ggplot(diamonds, aes(x=carat))`

`b=a+geom_histogram(bins=50)`

`b`

Draw the histogram of the variable *carat* with bins equal to 0.10

`a=ggplot(diamonds, aes(x=carat))`

`b=a+geom_histogram(binwidth=0.10)`

`b`

Draw the histogram (with densities and 50 bins) and the density plot (in blue) of the variable *carat*

```
hist(carat, 50, freq=F)
lines(density(carat), col="blue", lwd=2)
```

or

```
a=ggplot(diamonds, aes(carat))
b=a+geom_histogram(bins=50, aes(y = after_stat(density)))
c=b+geom_density(col="blue")
c
```

Draw a red and green histogram (with densities and 50 bins) and the density plot (in blue) of the variable *carat*

```
hist(carat, 50, freq=F, col="green")
lines(density(carat), col="blue", lwd=2)
```

or

```
a=ggplot(diamonds, aes(x=carat))
b=a+geom_histogram(bins=50, color="red", fill="green", aes(y = after_
stat(density)))
c=b+geom_density(col="blue")
c
```

Define the variable *price* from the dataset *diamonds*

```
price=diamonds$price
```

Draw the scatterplot between variables *carat* and *price*

```
plot(carat, price)
```

or

```
a=ggplot(diamonds,aes(x=carat, y=price))
b=a+geom_point()
b
```

Draw a smoothed scatterplot between variables *carat* and *price*

```
smoothScatter(carat, price)
```

or

```
a=ggplot(diamonds, aes(x=carat, y=price))
b=a+geom_bin2d(bins=250)
b
```

Compute the correlation between variables *carat* and *price*

```
cor(carat, price)
```

Run the correlation test to study between variables *cut* and *color*

Null hypothesis $H_0: r = 0$ (no correlation)

Alternative hypothesis $H_1: r \neq 0$ (correlation)

The decision of the test depends on the *p*-value.

If *p*-value < 0.05, there is an evidence against H_0 (we reject H_0).

If *p*-value ≥ 0.05 , there is an evidence in favor of H_0 (we do not reject H_0)

```
cor.test(carat, price)
```

Draw the scatterplot matrix of the variables *carat*, *price*, and *x* of the dataset *diamonds*

```
x=diamonds$x  
pairs(cbind(carat, price, x))
```

or

```
library(GGally)  
ggpairs(as.data.frame(cbind(carat, price, x)))
```

Data handling

Check the presence of missing values in a dataset
`anyNA(diamonds)`

If the output is FALSE, we have no missing value.

If the output is TRUE, we have one or more missing values, and we can identify them
`colSums(is.na(diamonds))`

If you want to delete the rows with NA (if any), we define a new dataset
`diamonds2=na.omit(diamonds)`

Check the presence of possible outliers for the variable *carat*
`boxplot(x)`

If there are possible outliers they can be listed using
`outliersx=boxplot(x)$out`

Regression analysis

Suppose we have imported the dataset *DatasetMarketing* reporting for 170 companies the following variables:

Nationality

Youtube advertising expenses

Facebook advertising expenses

Newspaper advertising expenses

Sales

The aim is the estimate of the best model to predict the sales of a company

First, check for the presence of missing values

`anyNA(DatasetMarketing)`

Given the answer (TRUE), we check how many missing values are present.

`colSums(is.na(DatasetMarketing))`

Given that there is just one missing value, we can omit the row containing it and define a new dataset

`DatasetMarketing2=na.omit(DatasetMarketing)`

Define the variables

```
nationality=DatasetMarketing2$nationality  
youtube=DatasetMarketing2$youtube  
facebook=DatasetMarketing2$facebook  
news=DatasetMarketing2$newspaper  
sales=DatasetMarketing2$sales
```

Then, we check for possible outliers and inaccuracies.

Next step is the estimate of the complete regression model

`mod0=lm(sales ~ nationality+youtube+facebook+news)`

Estimate the best model and display the output

```
modB=step(mod0)  
summary(modB)
```

Make a prediction with the following categories/values:

Nationality=US

Youtube expenses = 300

Facebook expenses = 100

Newspaper expenses = 100

```
newdata=data.frame(nationality="US", youtube=300, facebook=100, news=100)  
predict(modB, newdata)
```